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Report ID: March 19, 2002, 14:24:26 ; search time 151.3 seconds
(without alignments)
Memory: 64.192 Mbytes; update 5/25/01

Total hits: 10,114 on mask 1
Return scores: 0.5 - 9.86e-009.1
Sequences: 1,941 unique sequences
Orientation: FORWARD
Gap Penalty: Gapext: 1.0

Started:

361200 seqs, 113238999 residues

Total number of hits satisfying choice: Exact, etc.: 5
Minimum DB seq length: 10
Maximum DB seq length: 200000000

Post processing: Minimum Batch: 0%

Maximum Match: 100%

Listings: First 45 summaries

Database: Issued Patents NA *

Sequence 1, Appl

Sequence 2, Appl

Sequence 3, Appl

Sequence 4, Appl

Sequence 5, Appl

Sequence 6, Appl

Sequence 7, Appl

Sequence 8, Appl

Sequence 9, Appl

Sequence 10, Appl

Sequence 11, Appl

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